

IN THE SPECIFICATION

1. At page 4, lines 13-16, please substitute the following paragraph:

Figure 4A-4D Anatomic effects of drugs that destabilize microtubules on various xenografts (CaPan-1, HuCC-T1, SW480, and HT-29, respectively)

Photographs of mice bearing the indicated xenografts were treated with C. novyi-NT with HTI-286 and photographed 3 days later.

2. At page 6, lines 10-17, please substitute the following paragraph:

Figure 10A-10F Quantification of the effects of C. novyi-NT plus low doses of HTI-286 and L-NNA on various xenografts (HCT116, HuCC-T1, HT-29, SW480, B16, and DLD-1, respectively).

Animals (n > 6 for each arm) were treated with HTI-286 (1 mg/kg) + L-NNA (10 mg/kg) +/- C. novyi-NT. HTI-286 at this dose had no significant effects on tumor growth when administered without L-NNA, and vice versa. The p-values are based on comparisons between L-NNA + HTI-286 and L-NNA + HTI-286 + C. novyi-NT spores. The p-value is indicated only when the result was statistically significant.

3. At page 12-13, paragraph spanning, please substitute the following paragraph:

Mice bearing HCT116 xenografts were treated with C. novyi-NT spores with or without HTI-286 (1.7 mg/kg) as described above. Following euthanasia 24-48 hr later, tumors were

harvested, weighed, minced, and Dounce-homogenized. Genomic DNA was isolated using the Qiagen Genomic DNA Buffer Set and Genomic-tip 100/G. 25 ng of DNA was used in a PCR using the conditions described in (Vogelstein and Kinzler, 1999). The following primers specific for the *C. novyi*-NT phospholipase C gene were used: 5'- AAGATGGTACAGGA ACTCATTC (SEQ ID NO:1) and 5'-GCTTGTCGAAATACCATGTTGC (SEQ ID NO:2). No detectable primer dimers formed and no nonspecific amplification of mouse or xenograft genomic DNA occurred with these primers and PCR conditions. Real-time PCR was performed using an iCycler, and threshold cycle numbers were calculated using the iCycler Optical system interface software (Bio-Rad Lab, Hercules, CA). Averages of the threshold cycle number (Ct) of triplicate measurements per tumor were obtained, and three different tumors were studied for each treatment group. The results were expressed as the difference between the Ct of the treated tumor and the Ct of untreated controls, normalized to tumor weight.

4. Please add the sequence listing submitted with this paper to the end of the application.